

SEQUENCE LISTING

(I)

- APPLICANT. ELAZAR RABBANI (i) JANNIS G. STAVRIANOPOULOS JAMES J. DONEGAN DAKALLIU NORMAN E KELKER DEAN L. ENGELHARDT
- TITLE OF INVENTION: NOVEL PROPERTY EFFECTING AND/OR PROPERTY EXHIBITING COMPOSITIONS FOR (ii) THERAPEUTIC AND DIAGNOSTIC USE
- NUMBER OF SEQUENCES: 42 (iii)
- CORRESPONDENCE ADDRESS: (iv)
 - ADDRESSEE: ENZO THERAPEUTICS, INC. (A)
 - C/O ENZO BIOCHEM, INC. STREET: (B) 527 MADISON AVENUE,

9TH FLOOR

- **NEW YORK** CITY: (C)
- STATE: NY (D)
- USA COUNTRY: (E)
- 10022 ZIP: (F)
- COMPUTER READABLE FORM: (v)
 - 3.5" Micro Floppy Disk. 1.44 KB MEDIUM TYPE: (A) **STORAGE**
 - IBM PC/XT/AT, IBM PS/2 OR COMPATIBLES COMPUTER. (B)
 - OPERATING SYSTEM: PC-DOS (C)
 - MICROSOFT WORD ___ ASCII TEXT (DOS) SOFTWARE. (D)
- CURRENT APPLICATION DATA: (vi)
 - APPLICATION NUMBER: US 08/978,636 (A)
 - FILING DATE: 25-NOVEMBER-1997 (B)
 - Not Yet Known CLASSIFICATION: (C)
- ATTORNEY/AGENT INFORMATION: (viii)
 - NAME: FEDUS, RONALD C. (A)
 - **REGISTRATION NUMBER: 32,567** (B)
 - ENZ-53(D3) REFERENCE/DOCKET NUMBER: (C)
- TELECOMUNICATION INFORMATION: (ix)
 - TELEPHONE: (212) 583-0100 (A)
 - (212) 583-0150 TELEFAX: (B)
- INFORMATION FOR SEQ ID NO: 1 (2)
 - SEQUENCE CHARACTERISTICS: (i)
 - 20 amino acids LLNGTH: (A)
 - amino acid TYPE: (B)
 - single STRANDEDNESS: (C)
 - linear TOPOLOGY: (D)

MOLECULE TYPE: peptide (ii) HYPOTHETICAL: NO (iii) SEQ ID NO: 1: SEQUENCE DESCRIPTION (ix) Gly Phe Phe Gly Ala He Ala Gly Phe Leu Glu Gly Gly Trp Glu Gly Met He Ala Gly 20 INFORMATION FOR SEQ ID NO. 2: (3) SEQUENCE CHARACTERISTICS: (i) 20 base pairs LENGTH: (A) nucleic acid TYPE: (B) double STRANDEDNESS: (C) TOPOLOGY: linear (D) MOLECULE TYPE: DNA (genomic) (ii) HYPOTHETICAL: NO (iii) SEQ ID NO: 2: SEQUENCE DESCRIPTION: (ix) TGCTCTCTAA GGGTCTACTC 20 INFORMATION FOR SEQ ID NO. 3: (4) SEQUENCE CHARACTERISTICS: (i) 15 base pairs LENGTH: (A) nucleic acid TYPE. (B) double STRANDEDNESS: (C) linear TOPOLOGY: (D) MOLECULE TYPE: DNA (genomic) (ii) HYPOTHETICAL: NO (iii) SEQ ID NO: 3: SEQUENCE DESCRIPTION: (xi)CTCTAAGGTA AATAT 15 INFORMATION FOR SEQ ID NO. 4: (5) SEQUENCE CHARACTERISTICS: (i) 16 base pairs LENGTH: (A) nucleic acid TYPE:

(B)

(C)

(D)

STRANDEDNESS:

TOPOLOGY:

double

Iinear

MOLECULE TYPE: DNA (genomic) (ii) HYPOTHE FICAL: NO (iii) SEQ ID NO: 4: SEQUENCE DESCRIPTION: (ix) TGTATTTTAG ATTCAA 16 INFORMATION FOR SEQ ID NO. 5: (6) SEQUENCE CHARACTERISTICS: (i) 19 base pairs LENGTH: (A) nucleic acid TYPE (B) double STRANDEDNESS: (C) TOPOLOGY: linear (D) MOLECULE TYPE: DNA (genomic) (ii) HYPOTHETICAL: YES (iii) SEQ ID NO: 5: SEQUENCE DESCRIPTION: (ix) TGCTCTCTAA GGTAAATAT 19 INFORMATION FOR SEQ ID NO. 6. (7) SEQUENCE CHARACTERISTICS: (i) 19 base pairs LENGTEI: (A) nucleic acid TYPE: (B) double STRANDEDNESS: (C) TOPOLOGY: linear (D) MOLECULE TYPE: DNA (genomic) (ii) HYPOTHETICAL: YES (iii) 6: SEQ ID NO: SEQUENCE DESCRIPTION: (ix) IGTATTITAG GGTCTACTC 10 INFORMATION FOR SEQ ID NO 7: (8) SEQUENCE CHARACTERISTICS: (i) 19 base pairs LENGTH: (A)

nucleic acid

linear

single

TYPE:

STRANDEDNESS:

TOPOLOGY:

MOLECULE TYPE: mRNA

(B)

(C)

(D)

(ii)

- РА
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

UGCUCUCUAA GGUAAAUAU 19

- (9) INFORMATION FOR SEQ ID NO. 8:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

19 base pairs

- (B) TYPE:
- nucleic acid
- (C) STRANDEDNESS:
- : single
- (D) TOPOLOGY:
- Iinear
- (ii) MOLECULE TYPE: mRNA
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

UGUAUUUUAG GGUCUACUC 19

(10)INFORMATION FOR SEQ ID NO. 9:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH.

20 base pairs

- (B) TYPE:
- nucleic acid

linear

- (C) STRANDEDNESS:
- single
- (D) TOPOLOGY:
- •
- (ii) MOLECULE TYPE: mRNA
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

UGCUCUCUAA GGGUCUACUC 20

(11) INFORMATION FOR SEQ ID NO. 10

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 49 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (Λ) Description. desc = "oligonucleotide"
- (iii) HYPOTHETICAL: YES

(ix) SEQUENCE DESCRIPTION: SEQ ID NO: 10:

GGAATTCGTC TCGAGCTCTG ATCACCACCA TGGACACGAT TAACATCGC 49

(12)INFORMATION FOR SEQ ID NO. 11:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 55 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: /desc = "oligonucleotide"
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 11:

GACTAGTTGG TCTCGTCTCT TTTTTGGAGG AGTGTCGTTC TTAGCGATGT TAATC 55

(13) INFORMATION FOR SEQ ID NO. 12:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 46 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: 'desc = "oligonucleotide"
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 12:

GGAATTCGTC TCGGAGAAAG GTAAAATTCT CTGACATCGA ACTGGC

(14) INFORMATION FOR SEQ ID NO. 13:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs nucleic acid
 - (B) TYPE: nucleic acid (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: desc "oligonucleotide"

- ПҮРОТПЕТІСАL: YES (iii)
- 13: SEQ ID NO: SEQUENCE DESCRIPTION: (ix)

GACTAGTGGT CTCCCCTTAG AGAGCATGTC AGC 33

(15) INFORMATION FOR SEQ ID NO. 14:

- SEQUENCE CHARACTERISTICS: (i)
 - LENGTH: (A)
- 33 base pairs
- TYPE: (B)
- nucleic acid single
- STRANDEDNESS: (C)
 - TOPOLOGY: linear
- (D)
- MOLECULE TYPE: other nucleic acid (ii) (A) Description: /desc = "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- 14: SEQ ID NO: SEQUENCE DESCRIPTION: (ix)

GGAATTCGGT CTCGGGTCTA CTCGGTGGCG AGG 33

(16) INFORMATION FOR SEQ ID NO. 15:

- SEQUENCE CHARACTERISTICS: (i)
 - 27 base pairs LENGTH: (A)
 - nucleic acid (B) TYPE:
 - single STRANDEDNESS: (C)
 - linear TOPOLOGY: (D)
- MOLECULE TYPE: other nucleic acid (ii) (A) Description. /desc = "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- SEQ ID NO: 15: SEQUENCE DESCRIPTION: (ix)

GACTAGTCGT TACGCGAACG CAAAGTC

- INFORMATION FOR SEQ ID NO. 16: (17)
 - SEQUENCE CHARACTERISTICS: (i) 36 base pairs
 - LENGTII: (A)
 - nucleic acid TYPE: (B) STRANDEDNESS: single
 - (C) linear
 - TOPOLOGY: (D)
 - MOLECULE TYPE: other nucleic acid (ii) (A) Description: desc = "oligonucleotide"

- - HYPOTHETICAL: YES (iii)
 - SEQ ID NO: 16: SEQUENCE DESCRIPTION: (ix)

GGAATTCGTC TCTAAGGTAA ATATAAAATT TTTAAG 36

- INFORMATION FOR SEQ ID NO. 17: (18)
 - SEQUENCE CHARACTERISTICS: (i)
 - LENGTH: (A)
- 40 base pairs
- TYPE: (B)
- nucleic acid single
- STRANDEDNESS: (C)
- TOPOLOGY: linear (D)
- MOLECULE TYPE other nucleic acid (ii) (A) Description /desc = "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- 17: SEQUENCE DESCRIPTION: SEQ ID NO: (ix)

GACTAGTCGT CTCTGACCCT AAAATACACA AACAATTAGA 40

- INFORMATION FOR SEQ ID NO. 18: (19)
 - SEQUENCE CHARACTERISTICS: (i)
 - 92 base pairs LENGTH: (A)
 - nucleic acid (B) TYPE:
 - double STRANDEDNESS: (C)
 - TOPOLOGY: linear (D)
 - MOLECULE TYPE: DNA (genomic) (ii)
 - HYPOTHETICAL: YES (iii)
 - 18: SEQ ID NO: SEQUENCE DESCRIPTION: (ix)

GGAATTCGTC TCGAGCTCTG ATCACCACCA TGGACACGAT TAACATCGCT AAGAACGACA 60

CICCICCAAA AAAGAGACGA GACCAACTAG IC 92

- INFORMATION FOR SEQ ID NO. 19: (20)
 - SEQUENCE CHARACTERISTICS: (i)
 - 46 base pairs LENGTH. (A)
 - nucleic acid TYPE: (B)
 - STRANDEDNESS: double (C)
 - linear TOPOLOGY: (D)

- MOLECULE TYPE DNA (genomic) (ii)
- HYPOTHETICAL: YES (iii)
- 19: SEQUENCE DESCRIPTION: SEQ ID NO: (ix)

GGAATTCGTC TCGGAGAAAG GTAAAATTCT CTGACATCGA ACTGGC

INFORMATION FOR SEQ ID NO. 20: (21)

- SEQUENCE CHARACTERISTICS: (i)
 - LENGTH: (A)

106 base pairs

(B) TYPE: nucleic acid double

STRANDEDNESS: (C)

- linear TOPOLOGY: (D)
- MOLECULE TYPE: DNA (genomic) (ii)
- HYPOTHETICAL YES (iii)
- SEQUENCE DESCRIPTION: (ix)

SEQ ID NO:

20:

GGAATTCGTC TCGAGCTCTG ATCACCACCA TGGACACGAT TAACATCGCT AAGAACGACA

CTCCTCCAAA AAAGAGAAAG GTAAAATTCT CTGACATCGA ACTGGC 106

INFORMATION FOR SEQ ID NO. 21: (22)

- SEQUENCE CHARACTERISTICS: (i)
 - LENGTH: (A)

50 base pairs

TYPE: (B)

nucleic acid

- (C)
- double STRANDEDNESS:
- linear TOPOLOGY: (D)
- MOLECULE TYPE: DNA (genomic) (ii)
- HYPOTHETICAL: NO (iii)
- SEQUENCE DESCRIPTION: (ix)

SEQ ID NO:

21:

ATGGACACGA ITAACATCGC TAAGAACGAC TTCTCTGACA TCGAACIGGC 50

INFORMATION FOR SEQ ID NO. 22: (23)

- SEQUENCE CHARACTERISTICS: (i)
 - LENGTH: (A)

77 base pairs

TYPE: (B)

nucleic acid

STRANDEDNESS: (C)

double

TOPOLOGY: linear (D)

- MOLECULE TYPE: DNA (genomic) (ii)
- HYPOTHETICAL: YES (iii)
- 22: SEQUENCE DESCRIPTION: SEQ ID NO: (ix)

ATGGACACGA TTAACATCGC TAAGAACGAC ACTCCTCCAA AAAAGAGAAA 60 **GGTAAAATTC**

TCTGACATCG AACTGGC 77

- INFORMATION FOR SEQ ID NO. 23: (24)
 - SEQUENCE CHARACTERISTICS: (i)

LENGTH: (A)

69 base pairs

TYPE: (B)

nucleic acid

STRANDEDNESS: (C)

single

- TOPOLOGY: linear (D)
- MOLECULE TYPE: other nucleic acid (ii) (A) Description: /desc = "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- SEQ ID NO: 23: SEQUENCE DESCRIPTION: (ix)

GATCATTAGA CCAGATCTGA GCCTGGGAGC TCTCTGGCTA ACTAGGGAAC 60 **CCACTGCTTA**

AGCCTCAAG 69

- INFORMATION FOR SEQ ID NO. 24: (25)
 - SEQUENCE CHARACTERISTICS: (i)

LENGTH: (A)

69 base pairs

nucleic acid

TYPE: (B) STRANDEDNESS:

single

(C) linear TOPOLOGY: (D)

- MOLECULE TYPE: other nucleic acid (ii) (A) Description. desc "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- 24: SEQUENCE DESCRIPTION: SEQ ID NO: (ix)

GATCCTTGAG GCTTAAGCAG TGGGTTCCCT AGTTAGCCAG AGAGCTCCCA GGCTCAGATC

TGGTCTAAT 69

INFORMATION FOR SEQ ID NO. 25: (26)SEQUENCE CHARACTERISTICS: (i) 61 base pairs LENGTH: (A) nucleic acid (B) TYPE: single STRANDEDNESS: (C) TOPOLOGY: linear (D) MOLECULE TYPE: other nucleic acid (ii) (A) Description: 'desc = "oligonucleotide" HYPOTHETICAL: YES (iii) 25: SEQ ID NO: SEQUENCE DESCRIPTION: (ix) GATCACCTTA GGCTCTCCTA TGGCAGGAAG AAGCGGAGAC AGCGACGAAG ACCTCCTCAA G 61 INFORMATION FOR SEQ ID NO. 26: (27)SEQUENCE CHARACTERISTICS: (i) 61 base pairs LENGTH: (A) nucleic acid TYPE: (B) single STRANDEDNESS: (C) TOPOLOGY: linear (D) MOLECULE TYPE: other nucleic acid (ii) (A) Description: /desc = "oligonucleotide" HYPOTHETICAL: YES (iii) SEQ ID NO: 26: SEQUENCE DESCRIPTION: (ix) GATCCTTGAG GAGGTCTTCG TCGCTGTCTC CGCTTCTTCC TGCCATAGGA 60 GAGCCTAAGG T 6 E INFORMATION FOR SEQ ID NO. 27: (28)SEQUENCE CHARACTERISTICS: (i) 62 base pairs (A) LENGTH: nucleic acid (B) TYPE: single STRANDEDNESS: (C) TOPOLOGY: linear (D)

MOLECULE TYPE: other nucleic acid

(A) Description: desc = "oligonucleotide"

(ii)

- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 27:

GATCATAGTG AATAGAGTTA GGCAGGGATA CTCACCATTA TCGTTTCAGA CCCACCTCCC 60

 ΔG

62

- (29) INFORMATION FOR SEQ ID NO 28:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

62 base pairs

- (B) TYPE:
- nucleic acid SS: single
- (C) STRANDEDNESS:
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description 'desc "oligonucleotide"
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 28:

GATCCTGGGA GGTGGGTCTG AAACGATAAT GGTGAGTATC CCTGCCTAAC TCTATTCACT 60

ΑT

62

- (30) INFORMATION FOR SEQ ID NO. 29:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

30 base pairs

- A) ELNOT
- nucleic acid
- (B) TYPE:
- single
- (C) STRANDEDNESS:(D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description desc = "oligonucleotide"
- (iii) HYPOTHETICAL, YES
- (iv) ORIGINAL SOURCE: (A) ORGANISM: TER-1
- (x) SEQUENCE DESCRIPTION: SEQ ID NO: 29:

AATCTAGAGC TAACAAAGCC CGAAAGGAAG 30

INFORMATION FOR SEQ ID NO. 30: (31) SEQUENCE CHARACTERISTICS: (i) LENGTH:

- nucleic acid TYPE (B) single STRANDEDNESS: (C)
- TOPOLOGY: linear (D)
- MOLECULE TYPE: other nucleic acid (ii) (A) Description desc "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- 30: SEQUENCE DESCRIPTION: SEO ID NO: (ix)

TTCTGCAGAT ATAGTTCCTC CTTTCAGC 28

INFORMATION FOR SEQ ID NO. 31: (32)

- SEQUENCE CHARACTERISTICS: (i)
 - LENGTH: (A)

70 base pairs

28 base pairs

TYPE: (B)

nucleic acid single

STRANDEDNESS: (C)

- TOPOLOGY: linear (D)
- MOLECULE TYPE: other nucleic acid (ii) (A) Description: /desc = "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- SEQ ID NO: 31: SEQUENCE DESCRIPTION: (ix)

TCGAGCCATG GCTTAAGGAT CCGTACGTCC GGAGCTAGCG GGCCCATCGA 60 **TACTAGTTAA**

ATGCAGATCT 70

INFORMATION FOR SEQ ID NO. 32: (33)

SEQUENCE CHARACTERISTICS: (i)

LENGTH: (Λ)

70 base pairs

nucleic acid

(B) TYPI:

single

- STRANDEDNESS: (C)
- TOPOLOGY: linear (D)
- MOLECULE TYPE: other nucleic acid (ii) (A) Description: -desc -- "oligonucleotide"
- HYPOTHETICAL: YES (iii)
- 32: SEO ID NO: SEQUENCE DESCRIPTION: (ix)

CTAGAGATCT GCATTTAACT AGTATCGATG GGCCCGCTAG CTCCGGACGT ACGGATCCTT 60

AAGCCATGGC 70

(34) INFORMATION FOR SEQ ID NO. 33:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 29 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: 'desc = "oligonucleotide"
- (iii) EIYPOTHETICAE: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 33:

CATGAAATTA ATTCGACTCA CTATACGGA 29

(35) INFORMATION FOR SEQ ID NO. 34:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 29 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: 'desc = "oligonuclcotide"
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 34:

TTTAATTAAG CTGAGTGATA TGCCTCTAG 29

(36) INFORMATION FOR SEQ ID NO. 35:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 72 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description desc "oligonucleotide"

- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 35:

GATCCGGATT GAGGCTTAAG CAGTGGGTTC CCTAGTTAGC CAGAGAGCTC CCAGGCTCAG 60

ATCTGGTCTA AT 72

- (37) INFORMATION FOR SEQ ID NO. 36:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

72 base pairs

- (B) TYPE:
- nucleic acid
- (C) STRANDEDNESS:
- single
- (D) TOPOLOGY:
 - linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: /desc = "oligonucleotide"
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 36:

CCGGATTAGA CCAGATCTGA GCCTGGGAGC TCTCTGGCTA ACTAGGGAAC CCACTGCTTA 60

AGCCTCAATC CG 72

- (38) INFORMATION FOR SEQ ID NO. 37:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

66 base pairs

- (B) TYPE.
- nucleic acid
- (C) STRANDEDNESS:
- : single
- (D) TOPOLOGY:
- linear
- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: 'desc = "oligonucleotide"
- (iii) HYPOTHETICAL, YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 37:

GATCCGGACC TTGAGGAGGT CTTCGTCGCT GTCTCCGCTT CTTCCTGCCA TAGGAGAGCC 60

TAAGGT

66

(39) INFORMATION FOR SEQ ID NO. 38:

SEQUENCE CHARACTERISTICS: (i) 66 base pairs LENGTH: nucleic acid TYPE: (B) single STRANDEDNESS: (C) TOPOLOGY: linear (D) MOLECULE TYPE, other nucleic acid (ii) (A) Description: desc -- "oligonucleotide" HYPOTHETICAL: YES (iii) 38: SEQUENCE DESCRIPTION: SEQ ID NO: (ix) CCGGACCTTA GGCTCTCCTA TGGCAGGAAG AAGCGGAGAC AGCGACGAAG ACCTCCTCAA GGTCCG 66 INFORMATION FOR SEQ ID NO. 39: (40)SEQUENCE CHARACTERISTICS: (i) 65 base pairs LENGTU: (A) nucleic acid TYPE: (B) STRANDEDNESS: single (C) linear TOPOLOGY: (D) MOLECULE TYPE: other nucleic acid (ii) (A) Description: /desc = "oligonucleotide" HYPOTHETICAL: YES (iii) SEQ ID NO: 39: SEQUENCE DESCRIPTION: (ix) GATCCGGATG GGAGGTGGGT CTGAAACGAT AATGGTGAGT ATCCCTGCCT **AACTCTATTC ACTAT** 65 INFORMATION FOR SEQ ID NO 40: (41)SEQUENCE CHARACTERISTICS: (i) LENGTH: 65 base pairs (A) nucleic acid TYPE: (B)

STRANDEDNESS:

MOLECULE TYPE: other nucleic acid

(A) Description | desc = "oligonucleotide"

TOPOLOGY:

HYPOTHETICAL: YES

(C)

(D)

(ii)

(iii)

single

linear

(ix) SEQUENCE DESCRIPTION: SEQ ID NO: 40:

CCGGATAGTG AATAGAGTTA GGCAGGGATA CTCACCATTA TCGTTTCAGA CCCACCTCCC 60

ATCCG

65

- (42) INFORMATION FOR SEQ ID NO. 41:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH:

67 base pairs

(B) TYPE:

nucleic acid SS: single

(C) STRANDEDNESS:

(D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: other nucleic acid
 (A) Description: 'desc "oligonucleotide"
- (iii) HYPOTHFTICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 41:

GATCAGCATG CCTGCAGGTC GACTCTAGAC CCGGGTACCG AGCTCGCCCT ATAGTGAGTC 60

GTATTAT 67

- (43) INFORMATION FOR SEQ ID NO. 42:
 - (i) SEQUENCE CHARACTERISTICS:

(A) LENGTH:

67 base pairs

(B) TYPE:

nucleic acid

(C) STRANDEDNESS:

single

(D) TOPOLOGY: linear

- (ii) MOLECULE TYPE: other nucleic acid(A) Description | desc = "oligonucleotide"
- (iii) HYPOTHETICAL: YES
- (ix) SEQUENCE DESCRIPTION: SEQ ID NO: 42:

CCGGATAATA CGACTCACTA TAGGGCGAGC FCGGTACCCG GGTCTAGAGT CGACCTGCAG 60

GCATGCT

67